0500-0420

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/058,291

DATE: 02/10/2002
TIME: 15:09:10

#2.

Input Set : N:\Crf3\RULE60\10058291.txt
Output Set: N:\CRF3\02102002\J058291.raw

SEQUENCE LISTING

```
 4 (1) GENERAL INFORMATION:
 6        (i) APPLICANT: Hartley, James L.
 7                       Brasch, Michael A.
 9       (ii) TITLE OF INVENTION: Recombinational Cloning Using Engineered
10                                Recombination Sites
12      (iii) NUMBER OF SEQUENCES: 35
14       (iv) CORRESPONDENCE ADDRESS:
15            (A) ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C
16            (B) STREET: 1100 New York Ave., N. W. Suite 600
17            (C) CITY: Washington
18            (D) STATE: DC
19            (E) COUNTRY: USA
20            (F) ZIP: 20005-3934
22        (v) COMPUTER READABLE FORM:
23            (A) MEDIUM TYPE: Floppy disk
24            (B) COMPUTER: IBM PC compatible
25            (C) OPERATING SYSTEM: PC-DOS/MS-DOS
26            (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
28       (vi) CURRENT APPLICATION DATA:
C--> 29       (A) APPLICATION NUMBER: US/10/058,291
C--> 30       (B) FILING DATE: 30-Jan-2002
56            (C) CLASSIFICATION:
C--> 53     (vii) PRIOR APPLICATION DATA:
34            (A) APPLICATION NUMBER: 09/432,085
35            (B) FILING DATE: 1999-11-02
39            (A) APPLICATION NUMBER: 09/233,493
40            (B) FILING DATE: 20-JAN-1999
44            (A) APPLICATION NUMBER: 09/005,476
45            (B) FILING DATE: 12-JAN-1998
49            (A) APPLICATION NUMBER: 08/663,002
50            (B) FILING DATE: 07-JUN-1996
54            (A) APPLICATION NUMBER: 08/486,139
55            (B) FILING DATE: 07-JUN-1995
58       (ix) TELECOMMUNICATION INFORMATION:
59            (A) TELEPHONE: 202-371-2600
60            (B) TELEFAX: 202-371-2540
63 (2) INFORMATION FOR SEQ ID NO: 1:
65        (i) SEQUENCE CHARACTERISTICS:
66            (A) LENGTH: 25 base pairs
67            (B) TYPE: nucleic acid
68            (C) STRANDEDNESS: both
69            (D) TOPOLOGY: both
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ENTERED

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71      (ii) MOLECULE TYPE: cDNA
76      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
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78 RKYCWGCTTT YKTRTACNAA STSGB
80 (2) INFORMATION FOR SEQ ID NO: 2:
82          (i) SEQUENCE CHARACTERISTICS:
83              (A) LENGTH: 25 base pairs
84              (B) TYPE: nucleic acid
85              (C) STRANDEDNESS: both
86              (D) TOPOLOGY: both
88      (ii) MOLECULE TYPE: cDNA
93      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
                                                                    25
95 AGCCWGCTTT YKTRTACNAA CTSGB
97 (2) INFORMATION FOR SEQ ID NO: 3:
99          (i) SEQUENCE CHARACTERISTICS:
100             (A) LENGTH: 25 base pairs
101             (B) TYPE: nucleic acid
102             (C) STRANDEDNESS: both
103             (D) TOPOLOGY: both
105     (ii) MOLECULE TYPE: cDNA
110     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                    25
112 GTTCAGCTTT CKTRTACNAA CTSGB
114 (2) INFORMATION FOR SEQ ID NO: 4:
116         (i) SEQUENCE CHARACTERISTICS:
117             (A) LENGTH: 25 base pairs
118             (B) TYPE: nucleic acid
119             (C) STRANDEDNESS: both
120             (D) TOPOLOGY: both
122     (ii) MOLECULE TYPE: cDNA
127     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
                                                                    25
129 AGCCWGCTTT CKTRTACNAA GTSGB
131 (2) INFORMATION FOR SEQ ID NO: 5:
133         (i) SEQUENCE CHARACTERISTICS:
134             (A) LENGTH: 25 base pairs
135             (B) TYPE: nucleic acid
136             (C) STRANDEDNESS: both
137             (D) TOPOLOGY: both
139     (ii) MOLECULE TYPE: cDNA
144     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                    25
146 .GTTCAGCTTT YKTRTACNAA GTSGB
148 (2) INFORMATION FOR SEQ ID NO: 6:
150         (i) SEQUENCE CHARACTERISTICS:
151             (A) LENGTH: 25 base pairs
152             (B) TYPE: nucleic acid
153             (C) STRANDEDNESS: both
154             (D) TOPOLOGY: both
156     (ii) MOLECULE TYPE: cDNA
161     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
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163 AGCCTGCTTT TTTGTACAAA CTTGT
165 (2) INFORMATION FOR SEQ ID NO: 7:
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167        (i) SEQUENCE CHARACTERISTICS:
168            (A) LENGTH: 25 base pairs
169            (B) TYPE: nucleic acid
170            (C) STRANDEDNESS: both
171            (D) TOPOLOGY: both
173       (ii) MOLECULE TYPE: cDNA
178       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:                   25
180 AGCCTGCTTT CTTGTACAAA CTTGT
182 (2) INFORMATION FOR SEQ ID NO: 8:
184        (i) SEQUENCE CHARACTERISTICS:
185            (A) LENGTH: 25 base pairs
186            (B) TYPE: nucleic acid
187            (C) STRANDEDNESS: both
188            (D) TOPOLOGY: both
190       (ii) MOLECULE TYPE: cDNA
195       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:                   25
197 ACCCAGCTTT CTTGTACAAA CTTGT
199 (2) INFORMATION FOR SEQ ID NO: 9:
201        (i) SEQUENCE CHARACTERISTICS:
202            (A) LENGTH: 25 base pairs
203            (B) TYPE: nucleic acid
204            (C) STRANDEDNESS: both
205            (D) TOPOLOGY: both
207       (ii) MOLECULE TYPE: cDNA
212       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:                   25
214 GTTCAGCTTT TTTGTACAAA CTTGT
216 (2) INFORMATION FOR SEQ ID NO: 10:
218        (i) SEQUENCE CHARACTERISTICS:
219            (A) LENGTH: 25 base pairs
220            (B) TYPE: nucleic acid
221            (C) STRANDEDNESS: both
222            (D) TOPOLOGY: both
224       (ii) MOLECULE TYPE: cDNA
229       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:                  25
231 GTTCAGCTTT CTTGTACAAA CTTGT
233 (2) INFORMATION FOR SEQ ID NO: 11:
235        (i) SEQUENCE CHARACTERISTICS:
236            (A) LENGTH: 25 base pairs
237            (B) TYPE: nucleic acid
238            (C) STRANDEDNESS: both
239            (D) TOPOLOGY: both
241       (ii) MOLECULE TYPE: cDNA
246       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:                  25
248 GTTCAGCTTT CTTGTACAAA GTTGG
250 (2) INFORMATION FOR SEQ ID NO: 12:
252        (i) SEQUENCE CHARACTERISTICS:
253            (A) LENGTH: 25 base pairs
254            (B) TYPE: nucleic acid
255            (C) STRANDEDNESS: both
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2/10/02

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256          (D) TOPOLOGY: both
258     (ii) MOLECULE TYPE: cDNA
263     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:                        25
265 AGCCTGCTTT TTTGTACAAA GTTGG
267 (2) INFORMATION FOR SEQ ID NO: 13:
269      (i) SEQUENCE CHARACTERISTICS:
270          (A) LENGTH: 25 base pairs
271          (B) TYPE: nucleic acid
272          (C) STRANDEDNESS: both
273          (D) TOPOLOGY: both
275     (ii) MOLECULE TYPE: cDNA
280     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:                        25
282 AGCCTGCTTT CTTGTACAAA GTTGG
284 (2) INFORMATION FOR SEQ ID NO: 14:
286      (i) SEQUENCE CHARACTERISTICS:
287          (A) LENGTH: 25 base pairs
288          (B) TYPE: nucleic acid
289          (C) STRANDEDNESS: both
290          (D) TOPOLOGY: both
292     (ii) MOLECULE TYPE: cDNA
297     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:                        25
299 ACCCAGCTTT CTTGTACAAA GTTGG
301 (2) INFORMATION FOR SEQ ID NO: 15:
303      (i) SEQUENCE CHARACTERISTICS:
304          (A) LENGTH: 25 base pairs
305          (B) TYPE: nucleic acid
306          (C) STRANDEDNESS: both
307          (D) TOPOLOGY: both
309     (ii) MOLECULE TYPE: cDNA
314     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:                        25
316 GTTCAGCTTT TTTGTACAAA GTTGG
318 (2) INFORMATION FOR SEQ ID NO: 16:
320      (i) SEQUENCE CHARACTERISTICS:
321          (A) LENGTH: 25 base pairs
322          (B) TYPE: nucleic acid
323          (C) STRANDEDNESS: both
324          (D) TOPOLOGY: both
326     (ii) MOLECULE TYPE: cDNA
331     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:                        25
333 GTTCAGCTTT CTTGTACAAA GTTGG
335 (2) INFORMATION FOR SEQ ID NO: 17:
337      (i) SEQUENCE CHARACTERISTICS:
338          (A) LENGTH: 39 base pairs
339          (B) TYPE: nucleic acid
340          (C) STRANDEDNESS: both
341          (D) TOPOLOGY: both
343     (ii) MOLECULE TYPE: cDNA
348     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:                        39
350 CCACCACAAA CGCGTCCATG GAATTACACT TTAATTTAG
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352 (2) INFORMATION FOR SEQ ID NO: 18:
354      (i) SEQUENCE CHARACTERISTICS:
355          (A) LENGTH: 39 base pairs
356          (B) TYPE: nucleic acid
357          (C) STRANDEDNESS: both
358          (D) TOPOLOGY: both
360      (ii) MOLECULE TYPE: cDNA
365      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
367 CCACCACAAG TCGACGCATG CCGACAGCCT TCCAAATGT                39
369 (2) INFORMATION FOR SEQ ID NO: 19:
371      (i) SEQUENCE CHARACTERISTICS:
372          (A) LENGTH: 46 base pairs
373          (B) TYPE: nucleic acid
374          (C) STRANDEDNESS: both
375          (D) TOPOLOGY: both
377      (ii) MOLECULE TYPE: cDNA
382      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
384 GGCCGATTAC GATATCCCAA CGACCGAAAA CCTGTATTTT CAGGGT       46
386 (2) INFORMATION FOR SEQ ID NO: 20:
388      (i) SEQUENCE CHARACTERISTICS:
389          (A) LENGTH: 30 base pairs
390          (B) TYPE: nucleic acid
391          (C) STRANDEDNESS: both
392          (D) TOPOLOGY: both
394      (ii) MOLECULE TYPE: cDNA
399      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
401 CAGGTTTTCG GTCGTTGGGA TATCGTAATC                         30
403 (2) INFORMATION FOR SEQ ID NO: 21:
405      (i) SEQUENCE CHARACTERISTICS:
406          (A) LENGTH: 47 base pairs
407          (B) TYPE: nucleic acid
408          (C) STRANDEDNESS: both
409          (D) TOPOLOGY: both
411      (ii) MOLECULE TYPE: cDNA
416      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
418 GGCCAGATTA CGATATCCCA ACGACCGAAA ACCTGTATTT TCAGGGT      47
420 (2) INFORMATION FOR SEQ ID NO: 22:
422      (i) SEQUENCE CHARACTERISTICS:
423          (A) LENGTH: 31 base pairs
424          (B) TYPE: nucleic acid
425          (C) STRANDEDNESS: both
426          (D) TOPOLOGY: both
428      (ii) MOLECULE TYPE: cDNA
433      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
435 CAGGTTTTCG GTCGTTGGGA TATCGTAATC T                       31
437 (2) INFORMATION FOR SEQ ID NO: 23:
439      (i) SEQUENCE CHARACTERISTICS:
440          (A) LENGTH: 48 base pairs
441          (B) TYPE: nucleic acid
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VERIFICATION SUMMARY   DATE: 02/10/2002
PATENT APPLICATION: US/10/058,291   TIME: 15:09:11

Input Set : N:\Crf3\RULE60\10058291.txt
Output Set: N:\CRF3\02102002\J058291.raw

L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:33 M:220 C: Keyword misspelled or invalid format, [(vii) PRIOR APPLICATION DATA:]